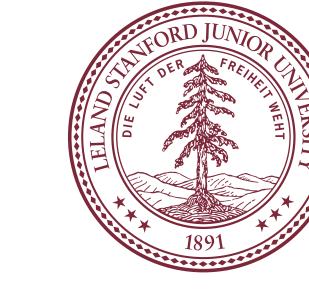


## Sources and Fate of Salmonella and Fecal Indicator Bacteria in an Urban Creek

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tound that San Pedro Creek in Pacifica had unusually high levels of *Salmonella*, relative to other nearby waterbodies. The present study investigated the spatial and temporal variability of Salmonella and Fecal Indicator Bacteria (FIB) in San Pedro Creek and used human and horse-specific Bacteroidales fecal markers to conduct microbial source tracking. Three surveys along San Pedro Creek spanning urban and forested areas were conducted. Salmonella was detected adjacent to and downstream of urban land cover, but not adjacent to forested land cover. No human or horsespecific *Bacteroidales* fecal markers were detected. Enterococci and *Salmonella* were positively correlated. Additionally, concentrations of *Salmonella* and enterococci were significantly higher during a survey conducted in wet weather compared to those conducted in dry weather. Two locations where Salmonella was consistently detected were sampled hourly for 25 hours and a mass-balance model was applied to determine field-relevant light and dark inactivation rates for Salmonella, enterococci, and Escherichia coli. Sunlight inactivation did not appear important in modulating concentrations of Salmonella, but was important in modulating both enterococci and *E. coli* concentrations. Dark inactivation was important for all three organisms. Inclusion of a zeroth order source term within the creek was needed to explain the dynamics of *E. coli*, suggesting this organism may grow within the creek itself.



-----Upstream

Modeled downstream

## Study Design

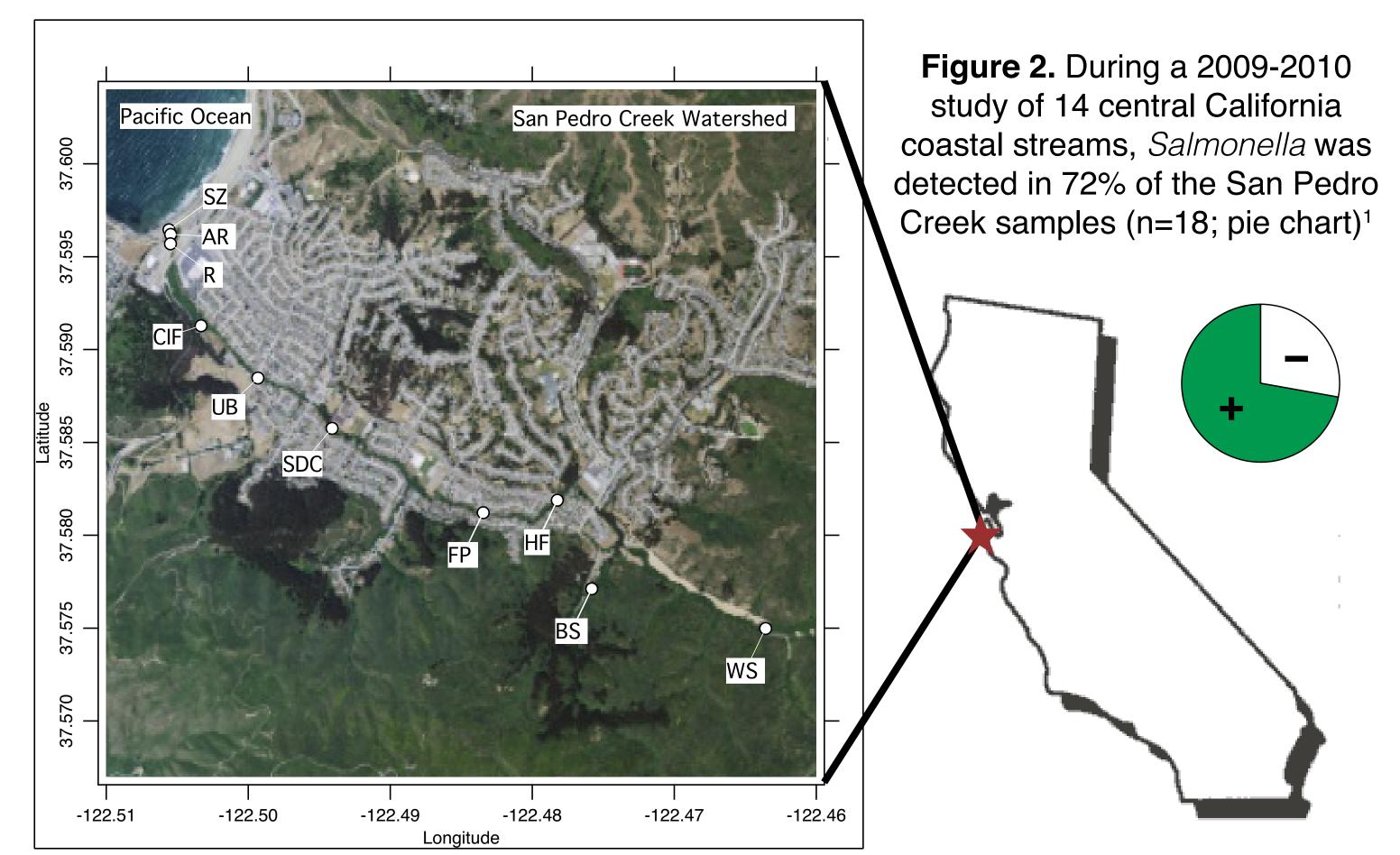


Figure 1. San Pedro Creek is a perennial creek draining a 20.7 km<sup>2</sup> watershed.

- May 24, 2010 4:00 h *E. coli* (CFU/100 ml)

## **Study Goals:**

1) Identify locations along San Pedro Creek where inputs of FIB and *Salmonella* occur

Microbiological Assays

enterococci (CFU/100 ml)

human-specific Bacteroidales (+/-)

horse-specific Bacteroidales (+/-)

- 2) Determine whether humans or horses contribute Salmonella or indicator bacteria to the creek using Bacteroidales host-specific fecal
- 3) Determine field-relevant dark and light inactivation rates of indicator bacteria and Salmonella within the creek using a mass-balance modeling approach using:  $v_{r} \frac{dC}{dt} = -kC + S$

C= concentration; x=distance along creek; v=longitudinal velocity; k=first order decay rate; S=source

3) L. K. Dick, A. E. Bernhard, T. J. Brodeur, J. W. Santo Domingo, J. M. Simpson, S. P. Walters and K. G. Field, Applied and Environment



# Salmonella 🔵 enterococci 🔘 E. coli 0 0 0 SZ AR R CIF UB SDC FP HF BS WS cator of *Salmonella* presence.

SZ AR R CIF UB SDC FP HF BS WS

SZ AR R CIF UB SDC FP HF BS WS

Figure 3. Spatial survey results for June (top)

Open circles indicate measurements below our

limit of detection (0.75 MPN/L).

sampling site (ocean to watershed)

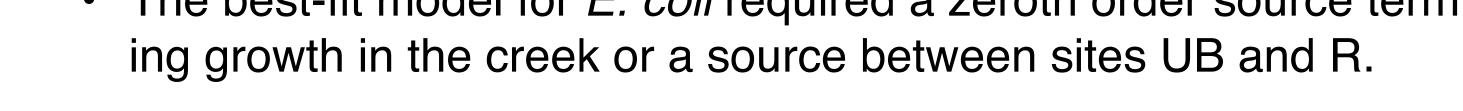
1000

- 1) Salmonella was consistently detected along the urban reach of San Pedro Creek and never in the undeveloped upper watershed (Figure 3).
- During dry weather Salmonella was found primarily between sites UB and R. • During wet weather, Salmonella was detected at all sites downstream of
- urban development, even in the surf zone. Salmonella and enterococci concentrations were correlated with each other, indicating that, in this watershed, the presence of enterococci is a good indi-
- 2) The human- and horse-specific markers were not detected in any samples suggesting that sources are non-human, non-equine, urban sources.

## **Temporal Results**

not *Salmonella* (Figure 4). The best-fit model for E. coli required a zeroth order source term suggest-

3) Results show evidence of sunlight inactivation for enterococci and *E. coli* but



## Table 1. Summary of best fit model parameters. k $(min^{-1})$ k $(min^{-1}l^{-1})$ Source (CFU/100 ml/min)

Organism	M <sub>dark</sub> (TTIIIT )	K <sub>sun</sub> (IIIIII I )	Jource (Or O) 100 IIIL/IIII
enterococci	4.4 x 10-3	1 x 10-4	0
E. coli	7.2 x 10-2	2.6 x 10-3	12.6
Salmonella	1 x 10-3	0	0

## **Human Health Risk**

Several serotypes of *Salmonella* known to cause human illness were found in San Pedro Creek including S. enterica Typhimirium, Enteritidis, and Heidelberg.4

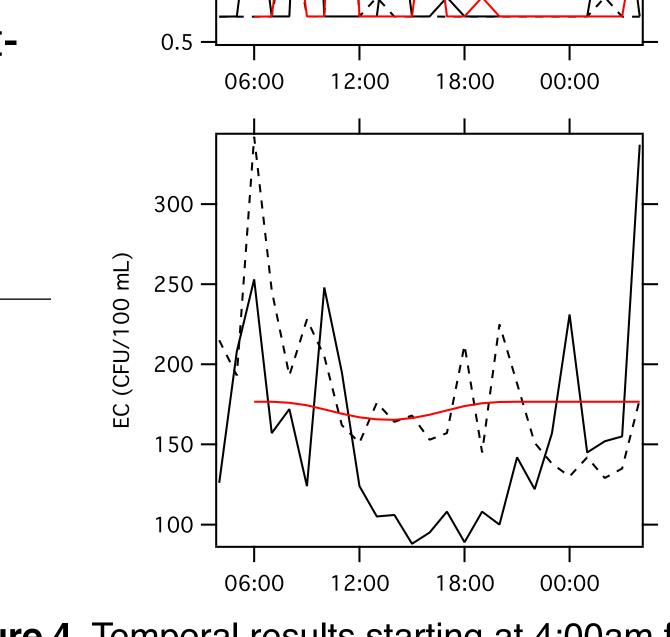


Figure 4. Temporal results starting at 4:00am for enterococci (top), *Salmonella* (middle) and *E. coli* (bottom) at the downstream site (UB) (black line), the upstream site (R) (dotted line), and the best-fit model (red line) for site R given site UB.

## Conclusions & Future Work

- While the health risk associated with the levels of Salmonella in San Pedro Creek are low, serotypes of Salmonella known to cause illness in humans were detected in the creek. Thus, exposure to the creek water, including the creek water discharged into the surf zone of Pacifica State Beach, should be avoided.
- Previous work in this watershed suggested that birds were a main source of *E. coli*, with additional contributions from dogs, horses, and humans.<sup>5</sup> Future work that applies avian, dog and human-specific source tracking methods will allow additional insight into the important sources of FIB and Salmonella in the creek.